



```

sequence extracted from NCBI backbone
REFERENCE
#note
A36007
#authors
Heller, R.A.; Song, K.; Onasch, M.A.; Fischer, W.H.; Chang, D.; Ringold, G.M.
#journal
Proc. Natl. Acad. Sci. U.S.A. (1990) 87:6151-6155
#title
Complementary DNA cloning of a receptor for tumor necrosis factor and demonstration of a shed form of the receptor.
#cross-references
MIMD:90349572
#accession
A36007
#status
Preliminary
#molecule_type
mRNA
#residues
116-140,'P',142-195,'R',197-362,'T',364-461 ##label HEL
#cross-references
GB:M35857
REFERENCE
#authors
Loetscher, H.; Schlaefer, E.J.; Iahn, H.W.; Pan, Y.C.E.; Lesslauer, W.; Brockhaus, M.
#journal
J. Biol. Chem. (1990) 265:1531-1536
#title
Purification and partial amino acid sequence analysis of two distinct tumor necrosis factor receptors from H160 cells.
#cross-references
MIMD:91050648
#accession
A23666
#status
Preliminary
#molecule_type
protein
#residues
23-40;65-69;136-141;300-306 ##label LOE
REFERENCE
#authors
Engelmann, H.; Novick, D.; Wallach, D.
#journal
J. Biol. Chem. (1990) 265:1531-1536
#title
Two tumor necrosis factor-binding proteins purified from human urine. Evidence for immunological cross-reactivity with cell surface tumor necrosis factor receptors.
#cross-references
MIMD:90110215
#accession
B35010
#status
Preliminary
#molecule_type
protein
#residues
27-31 ##label ENG
REFERENCE
#authors
Kuhner, P.; Kempf, O.; Wallach, D.
#journal
Gene (1994) 150:381-386
#title
Cloning, sequencing and partial functional characterization of the 5' region of the human p75 tumor necrosis factor receptor-encoding gene (TNF-R).
#cross-references
MIMD:95121934
#accession
I38094
#status
Preliminary; translated from GB/EMBL/DBJ
#molecule_type
DNA
#residues
1-37 ##label RES
#cross-references
EMBL:X80021; NID:g666044; CDS_PID:g825701
GENES
#note
GDB:TNFR2
#cross-references
GDB:125914
#map_position
1p36.2-1p36.2
26/3
CLASSIFICATION
#note
the list of introns is incomplete
#superfamily
tumor necrosis factor receptor type 2; NGF receptor repeat homology
duplication; receptor; transmembrane protein
KEYWORDS
#domain
signal sequence #status
predicted #label
SIG
#product
tumor necrosis factor receptor type 2 #status
experimental #label
MAY
#domain
NGF receptor repeat homology #label
NG1\
#domain
NGF receptor repeat homology #label
NG2\
#domain
NGF receptor repeat homology #label
NG3\
#domain
NGF receptor repeat homology #label
NG4\
#domain
transmembrane #status
predicted #label
TMN\
#domain
intracellular #status
predicted #label
INM\
#binding_site
carboxylate (asn) (covalent) #status
predicted
FEATURE
1-22
23-416
40-76
78-119
120-162
164-201
262-279
280-461
171,193
SUMMARY
#length
461 #molecular_weight
48291 #checksum
5724
13.1%; Score 398; DB 6; Length 461;
Query Match Best Local Similarity 43.08; Pred. 3.16e-47;
63; Conservative 19; Mismatches 55; Indels 7; Gaps 6

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[illegible]

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REFERENCE      A400254
#authors      Goodwin, R.G.; Anderson, D.; Jerzy, R.; Davis, T.; Brannan,
#journal      C.T.; Copeland, N.G.; Jenkins, N.A.; Smith, C.A.
#title        Mol. Cell. Biol. (1991) 11:3020-3026
#cross-references MBL:1246168
#accession    A40254
#molecule_type mRNA
#residues     1-474 ##label GOO
#cross-references GB:M60469
REFERENCE      S54816
#authors      Kisonergis, M.; Fellowes, R.; Feldmann, M.; Chernajovsky,
#submission   submitted to the EMBL Data Library, May 1995
#description   Characterization of the promoter region of the murine p75-TNFR
#receptor.
#accession    S54816
##molecule_type DNA
##residues     1-22 ##label K1S
#cross-references EMBL:X87128
CLASSIFICATION #superfamily tumor necrosis factor receptor type 2; NGF
#receptor repeat homology
FEATURE
1-22          #domain signal sequence #status predicted #label SIG\
23-474        #product tumor necrosis factor receptor type 2 #status
40-77         #domain NGF receptor repeat homology #label NG1\
79-120        #domain NGF receptor repeat homology #label NG2\
166-203       #domain NGF receptor repeat homology #label NG4
SUMMARY       #length 474 #molecular-weight 50319 #checksum 7767
Query Match   12.4%; Score 375; DB 6; Length 474;
Best Local Similarity 41.5%; Pred. No. 3,24e-43;
Matches 66; Conservative 21; Mismatches 61; Indels 11; Gaps 7;
Db 52 qmccakppgyvvhfnktsdtvcaadceasmytgvnqfritlscssscdtqyeiiaic 111
OY 38 QLLCDKCPPGYVYLTOKHTAKMTKVCAPCPDHYTDSMHTSDECTYCSPVCKELQYKQEC 97
Db 112 tkgnrycaceagaycaltkhsgscrgcmislscogpfgyastrapngnvlakaapgtf 171
OY 98 NRTINPVCCEKGRY--LEIEF--CLKH-R-S-CPPGFGVAGGPEKNTYCKRCPDGEF 150
Db 172 sdtstsvcrphricsi--laip--gnastdvacapes 206
OY 151 SNETSSKAPCKRHTNCYFGLLTQKGNATHDNCGNS 189
RESULT 4
ENTRY TITLE B60771 #type complete
ALTERNATE_NAMES B-cell activation protein CD40 precursor - human
ORGANISM B-cell surface antigen Bp50
DATE #format_name Homo sapiens #common_name man
03-Jun-1993 #sequence_revision 03-Feb-1994 #text_change
06-Sep-1996
ACCESSIONS S04460; A60771
REFERENCE S04460
#authors Stamenkovic, I.; Clark, E.A.; Seed, B.
#journal EMBO J. (1989) 8:1403-1410
#title A B-1-lymphocyte activation molecule related to the nerve
growth factor receptor and induced by cytokines in
carcinomas.
#cross-references MIMD:89356608
#accession S04460
#molecule_type mRNA
#residues 1-277 ##label STA
#cross-references EMBL:X60592
REFERENCE A60771
#authors Briesch Andersen, S.; Paulie, S.; Kono, H.; Nika, H.;
Aspenstrom, P.; Perlman, P.
#journal J. Immunol. (1989) 142:562-567

```

[illegible]

Qy 98 NTHNRVCECKEGRY-L--EIFECLKHRSRCPGFGVVOAGTPERNTVCKRCPDGFSSNET 154  
Db 156 slfeKcypwtscedknlewlqkqtsqtnvlg 187  
Qy 155 SSKAPCRKHTNCSVFGLLITOKGNATHDNICS 186

RESULT 6  
ENTRY A46476 #type complete  
TITLE CD40 - mouse  
ORGANISM #formal\_name Mus musculus #common\_name house mouse  
DATE 18-Jun-1993 #sequence\_revision 18-Nov-1994 #text\_change 18-Nov-1994

ACCESSIONS A46476  
REFERENCE A46476  
#authors Torres, R.M.; Clark, E.A.  
#journal J. Immunol. (1992) 148:620-626  
#title Differential increase of an alternatively polyadenylated mRNA species of murine CD40 upon B lymphocyte activation.  
#cross-references MUID:92105763  
#accession A46476  
##status preliminary  
##molecule\_type mRNA  
##residues 1-305 #label TOR  
##cross-references NCBIIN:75206; NCBIIP:75207  
#note Sequence extracted from NCBI backbone

SUMMARY #length 305 #molecular-weight 33617 #checksum 5203

Query Match 9.7%; Score 294; DB 14; Length 305;  
Best Local Similarity 38.8%; Pred. No. 2.21e-29;  
Matches 59; Conservative 20; Mismatches 65; Indels 8; Gaps 6;

Db 38 cdicppgsarllshcraletkqpcdsgefsagwrelchqhnrncp-qqrl-vkxeg 95  
Qy 41 CDKCPGGTYLKQHCYAKKTYCAPDPDHYTDSWHTSDECL-ly--CSPYCKELQYVQKQEC 97

Db 96 taesdtvtckegqhtskdcacaghpclpqlpfqvmematetdvtchpcpvgffsngs 155  
Qy 98 NTHNRVCECKEGRY-L--EIFECLKHRSRCPGFGVVOAGTPERNTVCKRCPDGFSSNET 154

Db 156 slfeKcypwtscedknlewlqkqtsqtnvlg 187  
Qy 155 SSKAPCRKHTNCSVFGLLITOKGNATHDNICS 186

RESULT 7  
ENTRY GQVZML #type complete  
TITLE T2 protein - myxoma virus (strain Lausanne)  
ORGANISM #formal\_name myxoma virus  
DATE 31-Dec-1992 #sequence\_revision 31-Dec-1992 #text\_change 26-Apr-1996

ACCESSIONS A40566  
REFERENCE A40566  
#authors Upton, C.; Macen, J.L.; Schreiber, M.; McFadden, G.  
#journal Virology (1991) 184:370-382  
#title Myxoma virus expresses a secreted protein with homology to the tumor necrosis factor receptor gene family that contributes to viral virulence.  
#cross-references MUID:91335768  
#accession A40566  
##molecule\_type DNA  
##residues 1-326 #label UPT  
##cross-references GB:M37976  
CLASSIFICATION #superfamily myxoma virus T2 protein; NGF receptor repeat homology

KEYWORDS glycoprotein

FEATURE 64-105 #domain NGF receptor repeat homology #label NG2\  
106-147 #domain NGF receptor repeat homology #label NG3\  
66,161,205,238 #binding\_site carbohydrate (asn) (covalent) #status predicted

SUMMARY #length 326 #molecular-weight 35208 #checksum 9255

Query Match 8.9%; Score 269; DB 2; Length 326;  
Best Local Similarity 33.8%; Pred. No. 3.14e-25;  
Matches 47; Conservative 25; Mismatches 58; Indels 9; Gaps 8;

Db 40 ctscppgyasrllogpsdvtcspcknetftastnhabacvscrgtqhlsegsedct 99  
Qy 41 CDKCPGGTYLKQHCYAKKTYCAPDPDHYTDSWHTSDECLYCSPVCKELQYVQKQECNRT 100

Db 100 rdryocdsagnycllkqgagccricapktkcpagysv-ghttrgdtlctkcpvtydsav 158  
Qy 101 HNRVCECKEGRY--LE-IEFC-L-KHRS-CPPGFGVVOAGTPERNTVCKRCPDGFSSNET 154

Db 159 sstetctssfnlysvetfnl 177  
Qy 155 SSKAPCRKHTNC-SV-FGL 171

RESULT 8  
ENTRY B43692 #type complete  
TITLE T2 protein - rabbit fibroma virus  
ORGANISM #formal\_name rabbit fibroma virus, Shope fibroma virus  
DATE 30-Sep-1993 #sequence\_revision 30-Sep-1993 #text\_change 26-Apr-1996

ACCESSIONS B43692  
REFERENCE B43692  
#authors Upton, C.; Delange, A.M.; McFadden, G.  
#journal Virology (1987) 160:20-30  
#title Tumorigenic poxviruses: genomic organization and DNA sequence of the telomeric region of the Shope fibroma virus genome.  
#accession B43692  
##status preliminary  
##molecule\_type DNA  
##residues 1-325 #label UPT  
##cross-references GB:M17433  
CLASSIFICATION #superfamily NGF receptor repeat homology

FEATURE 64-105 #domain NGF receptor repeat homology #label NG2\  
106-147 #domain NGF receptor repeat homology #label NG3\  
SUMMARY #length 325 #molecular-weight 35132 #checksum 4629

Query Match 8.6%; Score 260; DB 6; Length 325;  
Best Local Similarity 30.5%; Pred. No. 9.43e-24;  
Matches 51; Conservative 31; Mismatches 77; Indels 8; Gaps 5;

Db 40 cascbpgyasrllogpsntctvscpedgtftastnhabacvscrgtqhlsegsedct 99  
Qy 41 CDKCPGGTYLKQHCYAKKTYCAPDPDHYTDSWHTSDECLYCSPVCKELQYVQKQECNRT 100

Db 100 hdrvncstgnycllkqgngcrlicapgtkcpagysv-ghttrgdtlctkcpvtydsav 158  
Qy 101 HNRVCECKEGRY-L--EIE--FCIKHRSRCPGFGVVOAGTPERNTVCKRCPDGFSSNET 154

Db 159 sptercgtsfnlysvfnllypnetscett-aghnevktkfevtl 204  
Qy 155 SSKAPCRKHTNCSVFGLLITOKGNATHDNICSGNSESTOKGIDVTL 201

RESULT 9  
ENTRY I54182 #type complete  
ENTRY tumor necrosis factor receptor 2-related protein - human  
TITLE #formal\_name Homo sapiens #common\_name man  
ORGANISM #formal\_name Homo sapiens #common\_name man  
DATE 24-May-1996 #sequence\_revision 24-May-1996 #text\_change 24-May-1996

ACCESSIONS I54182  
REFERENCE I54182  
#authors Baens, M.; Chafanet, M.; Cassiman, J.J.; Van den Berghe, H.; Maynen, P.  
#journal Genomics (1993) 16:214-218  
#title Construction and evaluation of a hncDNA library of human 12p transcribed sequences derived from a somatic cell hybrid.  
#cross-references MUID:93252381  
#accession I54182  
##status preliminary; translated from GB/EMBL/DBJ

```
##molecule_type mRNA
##residues 1-435 ##label RES
##cross-references GB:U04270; NID:g339761; CDS_PID:g339762
SUMMARY #length 435 #molecular_weight 46709 #checksum 63

Query Match
  Best Local Similarity 32.3%; Pred. No. 9,436-24;
  Matches 52; Conservative 23; Mismatches 75; Indels 11; Gaps 6;

Db 52 epqhlccscppgtyvsakcsrltdvcatcaensynhmylltqlcrpdyv-mgl 110
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 34 ETSHOLLCDKCPGGTYLKHCHTAKMTVCAPCPDHYTDSMHTSDEC-LY--CSPVCKEL 90
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 111 eelap-cskrtktgcrcpymfcaawalectncellscdpppteaelkdyvkgmhcyp 169
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 91 OYVKOECHNHRVCECKEGRY----LEIEFLKHRSCPPGFGV-VQAGTPEKNTVCKR 144
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 170 ckaghfgntsparcqphtrcengqlveapgtagsdtic 210
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
SUMMARY #length 145 #checksum 185

RESULT 10
ENTRY D36858 #type complete
TITLE G4R protein - variola virus
ALTERNATE_NAMES B28R protein (COP)
ORGANISM #formal_name variola virus
DATE 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 15-Nov-1996

ACCESSIONS D36858; S46888; S35987
REFERENCE A36859
#authors Blinov, V.M.
#submission submitted to GenBank, November 1992
#description not shown.
#accession D36858
##status preliminary
##molecule_type DNA
##residues 1-349 ##label BLI
##cross-references GB:X69198
##experimental_source strain India-1967, ssp. major, isolate Ind3
REFERENCE S46888
#authors Kolyzhaylov, A.A.; Blinov, V.M.; Gyrtorov, V.V.; Pozdnyakov, S.G.; Chizhikov, V.E.; Frolov, I.V.; Totmenin, A.V.; Shenelkunov, S.N.; Sandakhchiev, L.S.
#submission submitted to the EMBL Data Library, April 1992
#description Nucleotide sequence analysis of the region of variola virus XhoI F O H P Q genome fragment.
#accession S46888
##status preliminary
##molecule_type DNA
##residues 1-349 ##label KOL
##cross-references EMBL:X67117
##experimental_source strain India-1967, isolate Ind3
CLASSIFICATION #superfamily NGF receptor repeat homology
FEATURE
  66-109 #domain NGF receptor repeat homology #label NG2\
  110-151 #domain NGF receptor repeat homology #label NG3
SUMMARY #length 349 #molecular_weight 38189 #checksum 2016

Query Match
  Best Local Similarity 32.0%; Pred. No. 2,206-19;
  Matches 34; Conservative 26; Mismatches 79; Indels 10; Gaps 7;

Db 1 mksvlylyllfscilindaaaytpmngkcdteykrhniclscppgyasrlcdsk 60
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 1 MNKLICALVFD-ISIKKTQETFPKYLHYD-E-TSHOLLCDKCPGGTYLKHCHTAK 57
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 61 tntgctpcsgtftsrnmhlpaciscngcrngqvetscnthmricecspgyycilkg 120
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 58 WKTVCAPCPDHYTDSMHTSDECILYCSPVCKELQYVKOCNRTNHRVCECKEGRY--LE- 114
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 121 ssgckacvsgtkcglygys-ghtsvgdvicspcgfgtyshtvsadkc 168
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :

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Oy 115 -IE--FCLKHRSCPPGFGVQAGTPEKNTVCKRCPPDGFSENTSSKAPC 160

RESULT 11
ENTRY S32385 #type fragment
TITLE gene G4R protein - variola virus (fragment)
ORGANISM #formal_name Variola virus
DATE 22-Nov-1993 #sequence_revision 22-Nov-1993 #text_change 22-Nov-1993

ACCESSIONS S32385
REFERENCE S32385
#authors Shchelkunov, S.N.; Blinov, V.M.; Sandakhchiev, L.S.
#journal FEBS Lett. (1993) 319:80-83
#title Genes of variola and vaccinia viruses necessary to overcome the host protective mechanisms.
#accession S32385
##status preliminary
##residues 1-138 ##label SHC
##cross-references EMBL:X69198
SUMMARY #length 138 #checksum 6036

Query Match
  Best Local Similarity 35.4%; Pred. No. 6,626-19;
  Matches 46; Conservative 19; Mismatches 58; Indels 7; Gaps 4;

Db 10 hnclscppgyasrlcdskntctpcgsgtftsrnmhlpaciscngcrngqvets 69
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 37 HOLLCDKCPGGTYLKHCHTAKMTVCAPCPDHYTDSMHTSDECILYCSPVCKELQYVKOE 96
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 70 cnthnriccspgyycilkgssgckacvsgtkcglygys-ghtsvgdvicspcgfgty 128
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 97 CNRTNHRVCECKEGRY--LE--IE--FCLKHRSCPPGFGVQAGTPEKNTVCKRCPPDGF 150
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 129 sltvsadkc 138
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 151 SNETSSKAPC 160
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :

RESULT 12
ENTRY I57826 #type complete
TITLE tumor necrosis factor receptor - mouse
ORGANISM #formal_name Mus musculus #common_name house mouse
DATE 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 02-Aug-1996

ACCESSIONS I57826
REFERENCE I57826
#authors Rothe, J.G.; Bluetmann, H.; Gentz, R.; Lesslauer, W.; Steinmetz, M.
#journal Mol. Immunol. (1993) 30:165-176
#title Genomic organization and promoter function of the murine tumor necrosis factor receptor beta gene.
#cross-references MUID:93156721
#accession I57826
##status preliminary; translated from GB/EMBL/DBJ
##molecule_type DNA
##residues 1-454 ##label RES
##cross-references GB:M76556; NID:g202100; CDS_PID:g202102
GENETICS
  #introns 13/3; 65/1; 108/1; 158/1; 184/2; 210/1; 248/1; 257/3; 353/1
  #exons gene name TNFR-2
SUMMARY #length 454 #molecular_weight 50030 #checksum 4267

Query Match
  Best Local Similarity 33.1%; Pred. No. 1,776-17;
  Matches 48; Conservative 21; Mismatches 65; Indels 11; Gaps 9;

Db 49 yvhsnmsicctkchkylysvdpcspgtrvcrecekglyftaagulylrglscctcke 108
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 31 YDEETSHOLLCDKCPGGTYLKHCHTAK-WKTVCAPCPDHYTDSMHTSDECILYCSPVCKE 89
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 109 msqvelspcgadkdvscgkngfgylysethfgvndspctng-tvllpckeqntvce 167
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Oy 90 LQYKA-QECNRTNHRVCECKEG--RYL-ETEF-CLKHRSCPPGFGVQAGTPEKNTVCK 143
   | : : : : : : : : : : : : : : : : : : : : : : : : : : : :

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Job time : 99 secs.

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